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Fig. 1

SEQ ID NO: 1

GGCGGCCGCTGCACTCCGCGCAGGCTGCGCCGCTGCACTCCGCGCAG GCTGTGGACGTGGCCTCGGCCTCCAACTTCCGGGCCTTTGAGCTGCTG CACTTGCACCTGGACCTGCGGGCTGAGTTCGGGCCTCCAGGGCCCGGC GCAGGGAGCCGGGGCTGAGCGCACCGCGGTCCTGGACCTGCGCTGC CTGGAGCCCGAGGCGCCGAGCTGCGCTGGACTCGCACCCGTGC CTGGAGGTGACGCGGCGCGCGCTGCGGCGGGAGCGGCCCGGCTCGGAG GAGCCGCCTGCGGAGCCCGTGAGCTTCTACACGCAGCCCTTCTCGCAC TATGGCCAGGCCCTGTGCGTGTCCTTCCCGCAGCCCTGCCGCGCCGCC GTTTGCTGGGTGCTCCCGAGCAGACAGCAGGAAAGAAGAAGCCCTTC GTGTACACCCAGGGCCAGGCTGTCCTAAACCGGGCCTTCTTCCCTTGC TTCGACACGCCTGCTGTTAAATACAAGTATTCAGCTCTTATTGAGGTC CCAGATGGCTTCACAGCTGTGATGAGTGCAAGACCTGGGGAGAAGAGA GGTCCAAATAAGTTCTTCTTCCAGATGTGTCAGCCCATCCCCTCTAT CTGATAGCTTTGGCCATCGGAGATCTGGTTTCGGCTGAAGTTGGACCC AGGAGCCGGGTGTGGGCTGAGCCCTGCCTGATTGATGCTGCCAAGGAG TACAACGGGGTGATAGAAGATTTTTGGCAACAGGAGAGAAGCTTTTT GGACCTTATGTTTGGGGAAGGTATGACTTGCTCTTCATGCCACCGTCC TTTCCATTTGGAGGAATGGAGAACCCTTGTCTGACCTTTGTCACCCCC TGCCTGCTAGCTGGGGACCGCTCCTTGGCAGATGTCATCATCCATGAG ATCTCCCACAGTTGGTTTGGGAACCTGGTCACCAACGCCAACTGGGGT GAATTCTGGCTCAATGAAGGTTTCACCATGTACGCCCAGAGGAGGATC TCCACCATCCTCTTTGGCGCTGCGTACACCTGCTTGGAGGCTGCAACG GGGCGGGCTCTGCTGCGTCAACACATGGACATCACTGGAGAGGAAAAC CCACTCAACAAGCTCCGCGTGAAGATTGAACCAGGCGTTGACCCGGAC GACACCTATAATGAGACCCCCTACGAGAAAGGTTTCTGCTTTGTCTCA

TACCTGGCCCACTTGGTGGGTGATCAGGATCAGTTTGACAGTTTTCTC AAGGCCTATGTGCATGAATTCAAATTCCGAAGCATCTTAGCCGATGAC TTTCTGGACTTCTACTTGGAATATTTCCCTGAGCTTAAGAAAAAGAGA GTGGATATCATTCCAGGTTTTGAGTTTGATCGATGGCTGAATACCCCC AAGCCTGCTGAAGAGCTAGCCCAACTGTGGGCAGCCGAGGAGCTGGAC ATGAAGGCCATTGAAGCCGTGGCCATCTCTCCCTGGAAGACCTACCAG CTGGTCTACTTCCTGGATAAGATCCTCCAGAAATCCCCTCTCCCT GGGAATGTGAAAAACTTGGAGACACATACCCAAGTATCTCAAATGCC CGGAATGCAGAGCTCCGGCTGCGATGGGGCCAAATCGTCCTTAAGAAC GACCACCAGGAAGATTTCTGGAAAGTGAAGGAGTTCCTGCATAACCAG GGGAAGCAGAAGTATACACTTCCGCTGTACCACGCAATGATGGGTGGC AGTGAGGTGGCCAGACCCTCGCCAAGGAGACTTTTGCATCCACCGCC TCCCAGCTCCACAGCAATGTTGTCAACTATGTCCAGCAGATCGTGGCA CCCAAGGGCAGTTAGAGGCTCGTGTGCATGGCCCCTGCCTCTTCAGGC TCTCCAGGCTTTCAGAATAATTGTTTGTTCCCAAATTCCTGTTCCCTG ATCAACTTCCTGGAGTTTATATCCCCTCAGGATAATCTATTCTCTAGC TCTATAGCCCACTGAGCCCCGAGACAGAGAACCTGCCCACAGCTCTCC CCGCTACAGGCTGCAGGCACGCAGGGCAGCGGGTATTCTCCTCCCACC TAAGTCTCTGGGAAGAAGTGGAGAGGACTGATGCTCTTCTTTTTCTC TTTCTGTCCTTTTTCTTGCTGATTTATGCAAAGGGCTGGCATTCTGAT GCTTTTCAGGTTTAATCCTTATTTTAATAAAGTTTTCAAGCAAAAAAA

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Fig. 2

SEQ ID NO: 2

MPPSFPFGGMENPCLTFVTPCLLAGDRSLADVIIHEISHSWFGNLVTN
ANWGEFWLNEGFTMYAQRRISTILFGAAYTCLEAATGRALLRQHMDIT
GEENPLNKLRVKIEPGVDPDDTYNETPYEKGFCFVSYLAHLVGDQDQF
DSFLKAYVHEFKFRSILADDFLDFYLEYFPELKKKRVDIIPGFEFDRW
LNTPGWPPYLPDLSPGDSLMKPAEELAQLWAAEELDMKAIEAVAISPW
KTYQLVYFLDKILQKSPLPPGNVKKLGDTYPSISNARNAELRLRWGQI
VLKNDHQEDFWKVKEFLHNQGKQKYTLPLYHAMMGGSEVAQTLAKETF
ASTASQLHSNVVNYVQQIVAPKGS

Fig. 3

SEQ ID NO: 3

5'-GAAGATTGAACCAGGCGTTGA-3'

Fig. 4

SEQ ID NO: 4

5'-GGTATGAGACAAAGCAGAAACCTTT-3'

Fig. 5

SEQ ID NO: 5

5'-CCGGACGACACCTATAATGAGACCCCCTAC-3'